

SEQUENCE LISTING

SEQ ID NO: 1

LENGTH: 1447

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION:

GAATTCGGG AAGGAATAAT TATTATTAGC AATTATTAGC GATCAATAAT CTTGATCACA TT 62

10 ATG GCA AGC ACT ATT AAG GAA GCA TTA TCA GIG GIG AGT GAA GAC CAG 110

TCC TTG TTT GAG TGT GGC TAC GGA TCG CCC CAC CTT GCA AAG ACA GAA 158

ATG ACA GCC TCC TCT TCC AGT GAA TAT GGG CAA ACA TCA AAG ATG AGC 206

CCG CGC GGT CCC CAG CAG GAC TGG TTA TCA CAG CCC CCG GCC AGA GGT 254

ACC ATT AAG ATG GAG TGT AAC CCA AAC CAG GGT AAT GGG TCA AGG AAT 302

15 TCA OCT GAT GAC TGC AGC GIG GCA AAA GGA GGG AAA ATG GGT AGC AGT 350

TCA GAC AAT GGT GGG ATG AAC TAT GGA AGC TAC ATG GAA GAG AAG CAT 398

ATT CCG OCT CCA AAT ATG ACA ACC AAT GAA CGA AGA GGT ATT GIG CCA 446

GCA GAT OCT ACG TTA TGG AGC ACA GAC CAT GTA CCG CAG TGG CTG GAG 494

TGG GCA GIG AAG GAG TAT GGT CTT CCA GAC GIG GAC ATC TTG TTG TTC 542

20 CAG AAC ATT GAT GGG AAA GAG TTG TGT AAA ATG ACC AAA GAT GAC TTC 590

CAG AGA CTC ACG CCG AGC TAT AAC GCA GAT ATC CTC CTG TCA CAC CTA 638

CAC TAC CTC AGA GAG AGA GGA GGC ACT TTT ATT TTT CCA AAT ACA TCA 686

GTT TAC CCA GAA GCA ACG CAA AGA ATA ACA ACA AGG CCA GAT TTA OCT 734

TAT GAG CAA GCG AGG AGA TCA GCG TGG ACG AGT CAC AGC CAT CCC ACT 782

25 CAG TCA AAA GCT ACC CAA CCA TCA TCT TCA ACA GIG CCC AAA ACA GAA 830

GAC CAG CGT CCT CAG TTA GAT OCT TAT CAG ATT CTT GGA CCG ACC AGC 878

AGC CGT CTT GCA AAT CCA GGG AGT GGG CAG ATA CAG CTA TGG CAG TTC 926
 CTA CTG GAG CTT CTG TCG GAC ACC TOC AAC TOC AAC TGC ATC ACC TGG 974
 GAG GGC ACA AAT GGG GAG TTC AAG ATG ACA GAC OCT GAT GAA GIG GCT 1022
 CGG CGT TGG GGA GAG AGG AAA AGC AAA OCT AAC ATG AAC TAT GAC AAA 1070
 CTC AGC CGT GCA CTT CGC TAC TAC TAT GAC AAA AAT ATT ATG ACT AAA 1118
 GGT CAT GGT AAA CGC TAT GGC TAC AAA TTT GAT TTC CAC GGA ATC GCT 1166
 CAG GGC CTC CAG OCT CAC OCT CCA GAA TCA TOC ATG TAC AAA TAC CCA 1214
 TCA GAC CTC CCC TAC ATG AGT TOC TAC CAT GCA CAC CCC CAG AAG ATG 1262
 AAC TTT GTA GCT CCC CAT CCC OCT GCT TTG CCC GTA ACC TCA TOC AGC 1310
 TTT TTT GCT GGC OCT AAT CCA TAC TGG AAT TCA CCA ACT GGA GGC ATC 1358
 TAC CCC AAT ACC AGG CTG CCA GCT GCT CAT ATG OCT TOC CAT CTT GGC 1406
 ACC TAC TAC TAA GIG GGGAAAGAAA GAAAGCGCCA AGAAAA 1447

SEQ ID NO: 2

LENGTH: 451

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION:

Met Ala Ser Thr Ile Lys Glu Ala Leu Ser Val Val Ser Glu Asp Gln 16
 Ser Leu Phe Glu Cys Ala Tyr Gly Ser Pro His Leu Ala Lys Thr Glu 32
 Met Thr Ala Ser Ser Ser Ser Glu Tyr Gly Gln Thr Ser Lys Met Ser 48
 Pro Arg Val Pro Gln Gln Asp Trp Leu Ser Gln Pro Pro Ala Arg Val 64

Thr Ile Lys Met Glu Cys Asn Pro Asn Gln Val Asn Gly Ser Arg Asn 80
 Ser Pro Asp Asp Cys Ser Val Ala Lys Gly Gly Lys Met Val Ser Ser 96
 Ser Asp Asn Val Gly Met Asn Tyr Gly Ser Tyr Met Glu Glu Lys His 112
 Ile Pro Pro Pro Asn Met Thr Thr Asn Glu Arg Arg Val Ile Val Pro 128
 Ala Asp Pro Thr Leu Trp Ser Thr Asp His Val Arg Gln Trp Leu Glu 144
 Trp Ala Val Lys Glu Tyr Gly Leu Pro Asp Val Asp Ile Leu Leu Phe 160
 Gln Asn Ile Asp Gly Lys Glu Leu Cys Lys Met Thr Lys Asp Asp Phe 176
 Gln Arg Leu Thr Pro Ser Tyr Asn Ala Asp Ile Leu Leu Ser His Leu 192
 His Tyr Leu Arg Glu Arg Gly Ala Thr Phe Ile Phe Pro Asn Thr Ser 208
 Val Tyr Pro Glu Ala Thr Gln Arg Ile Thr Thr Arg Pro Asp Leu Pro 224
 Tyr Glu Gln Ala Arg Arg Ser Ala Trp Thr Ser His Ser His Pro Thr 240
 Gln Ser Lys Ala Thr Gln Pro Ser Ser Ser Thr Val Pro Lys Thr Glu 256
 Asp Gln Arg Pro Gln Leu Asp Pro Tyr Gln Ile Leu Gly Pro Thr Ser 272
 Ser Arg Leu Ala Asn Pro Gly Ser Gly Gln Ile Gln Leu Trp Gln Phe 288
 Leu Leu Glu Leu Leu Ser Asp Ser Ser Asn Ser Asn Cys Ile Thr Trp 304
 Glu Gly Thr Asn Gly Glu Phe Lys Met Thr Asp Pro Asp Glu Val Ala 320
 Arg Arg Trp Gly Glu Arg Lys Ser Lys Pro Asn Met Asn Tyr Asp Lys 336
 Leu Ser Arg Ala Leu Arg Tyr Tyr Tyr Asp Lys Asn Ile Met Thr Lys 352
 Val His Gly Lys Arg Tyr Ala Tyr Lys Phe Asp Phe His Gly Ile Ala 368
 Gln Ala Leu Gln Pro His Pro Pro Glu Ser Ser Met Tyr Lys Tyr Pro 384
 Ser Asp Leu Pro Tyr Met Ser Ser Tyr His Ala His Pro Gln Lys Met 400
 Asn Phe Val Ala Pro His Pro Pro Ala Leu Pro Val Thr Ser Ser Ser 416
 Phe Phe Ala Ala Pro Asn Pro Tyr Trp Asn Ser Pro Thr Gly Gly Ile 432
 Tyr Pro Asn Thr Arg Leu Pro Ala Ala His Met Pro Ser His Leu Gly 448
 Thr Tyr Tyr 451

SEQ ID NO: 3

LENGTH: 1528

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION:

10 GAATTCGGG AACGAATAAT TATTATTAGC AATTATTAGC GATCAATAAT CTTGATCACA TT 62

ATG GCA AGC ACT ATT AAG GAA GCA TTA TCA GTG GTG AGT GAA GAC CAG 110

TCC TTG TTT GAG TGT GCC TAC GGA TCG CCC CAC CTT GCA AAG ACA GAA 158

ATG ACA GGC TCC TCT TCC AGT GAA TAT GGG CAA ACA TCA AAG ATG AGC 206

CCG CGC GGT CCC CAG CAG GAC TGG TTA TCA CAG CCC CCG GGC AGA GGT 254

15 ACC ATT AAG ATG GAG TGT AAC CCA AAC CAG GGT AAT GGG TCA AGG AAT 302

TCA OCT GAT GAC TGC AGC GTG GCA AAA GGA GGG AAA ATG GGT AGC AGT 350

TCA GAC AAT GGT GGG ATG AAC TAT GGA AGC TAC ATG GAA GAG AAG CAT 398

ATT CCG OCT CCA AAT ATG ACA ACC AAT GAA CGA AGA GGT ATT GTG CCA 446

GCA GAT OCT ACG TTA TGG AGC ACA GAC CAT GTA CCG CAG TGG CTG GAG 494

20 TGG GCA GTG AAG GAG TAT GGT CTT CCA GAC GTG GAC ATC TTG TTG TTC 542

CAG AAC ATT GAT GGG AAA GAG TTG TGT AAA ATG ACC AAA GAT GAC TTC 590

CAG AGA CTC ACG CCG AGC TAT AAC GCA GAT ATC CTC CTG TCA CAC CTA 638

CAC TAC CTC AGA GAG ACT OCT CTT CCA CAT TTG ACT TCA GAT GAT GGT 686

GAT AAG GGC TTA CAA AAC TCT CCA CCG TTA ATG CAT GCT AGA AAC ACA 734

25 GGA GGA GGC ACT TTT ATT TTT CCA AAT ACA TCA GTT TAC CCA GAA GCA 782

ACG CAA AGA ATA ACA ACA AGG CCA GAT TTA OCT TAT GAG CAA GCG AGG 830

AGA TCA GCG TGG ACG AGT CAC AGC CAT CCC ACT CAG TCA AAA GCT ACC 878
 CAA OCA TCA TCT TCA ACA GIG CCC AAA ACA GAA GAC CAG CGT OCT CAG 926
 TTA GAT OCT TAT CAG ATT CTT GGA CCG ACC AGC AGC CGT CTT GCA AAT 974
 OCA GGG AGT GGG CAG ATA CAG CTA TGG CAG TTC CTA CTG GAG CTT CTG 1022
 TCG GAC AGC TOC AAC TOC AAC TGC ATC ACC TGG GAG GGC ACA AAT GGG 1070
 GAG TTC AAG ATG ACA GAC OCT GAT GAA GIG GCT CCG CGT TGG GGA GAG 1118
 AGG AAA AGC AAA OCT AAC ATG AAC TAT GAC AAA CTC AGC CGT GCA CTT 1166
 CGC TAC TAC TAT GAC AAA AAT ATT ATG ACT AAA GAT CAT GGT AAA CGC 1214
 TAT GCC TAC AAA TTT GAT TTC CAC GGA ATC GCT CAG GCC CTC CAG OCT 1262
 CAC OCT OCA GAA TCA TOC ATG TAC AAA TAC OCA TCA GAC CTC CCC TAC 1310
 ATG AGT TOC TAC CAT GCA CAC CCC CAG AAG ATG AAC TTT GTA GCT CCC 1358
 CAT CCC OCT GCT TTG CCC GTA ACC TCA TOC AGC TTT TTT GCT GCC OCT 1406
 AAT OCA TAC TGG AAT TCA OCA ACT GGA GGC ATC TAC CCC AAT ACC AGG 1454
 CTG OCA GCT GCT CAT ATG OCT TOC CAT CTT GGC ACC TAC TAC TAA GIG 1502
 GGGAAAGAAA GAAAGCGCCA AGAAAA 1528

SEQ ID NO: 4

LENGTH: 478

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION:

Met Ala Ser Thr Ile Lys Glu Ala Leu Ser Val Val Ser Glu Asp Gln 16

Ser Leu Phe Glu Cys Ala Tyr Gly Ser Pro His Leu Ala Lys Thr Glu 32
 Met Thr Ala Ser Ser Ser Ser Glu Tyr Gly Gln Thr Ser Lys Met Ser 48
 Pro Arg Val Pro Gln Gln Asp Trp Leu Ser Gln Pro Pro Ala Arg Val 64
 Thr Ile Lys Met Glu Cys Asn Pro Asn Gln Val Asn Gly Ser Arg Asn 80
 Ser Pro Asp Asp Cys Ser Val Ala Lys Gly Gly Lys Met Val Ser Ser 96
 Ser Asp Asn Val Gly Met Asn Tyr Gly Ser Tyr Met Glu Glu Lys His 112
 Ile Pro Pro Pro Asn Met Thr Thr Asn Glu Arg Arg Val Ile Val Pro 128
 Ala Asp Pro Thr Leu Trp Ser Thr Asp His Val Arg Gln Trp Leu Glu 144
 Trp Ala Val Lys Glu Tyr Gly Leu Pro Asp Val Asp Ile Leu Leu Phe 160
 Gln Asn Ile Asp Gly Lys Glu Leu Cys Lys Met Thr Lys Asp Asp Phe 176
 Gln Arg Leu Thr Pro Ser Tyr Asn Ala Asp Ile Leu Leu Ser His Leu 192
 His Tyr Leu Arg Glu Thr Pro Leu Pro His Leu Thr Ser Asp Asp Val 208
 Asp Lys Ala Leu Gln Asn Ser Pro Arg Leu Met His Ala Arg Asn Thr 224
 Gly Gly Ala Thr Phe Ile Phe Pro Asn Thr Ser Val Tyr Pro Glu Ala 240
 Thr Gln Arg Ile Thr Thr Arg Pro Asp Leu Pro Tyr Glu Gln Ala Arg 256
 Arg Ser Ala Trp Thr Ser His Ser His Pro Thr Gln Ser Lys Ala Thr 272
 Gln Pro Ser Ser Ser Thr Val Pro Lys Thr Glu Asp Gln Arg Pro Gln 288
 Leu Asp Pro Tyr Gln Ile Leu Gly Pro Thr Ser Ser Arg Leu Ala Asn 304
 Pro Gly Ser Gly Gln Ile Gln Leu Trp Gln Phe Leu Leu Glu Leu Leu 320
 Ser Asp Ser Ser Asn Ser Asn Cys Ile Thr Trp Glu Gly Thr Asn Gly 336
 Glu Phe Lys Met Thr Asp Pro Asp Glu Val Ala Arg Arg Trp Gly Glu 352
 Arg Lys Ser Lys Pro Asn Met Asn Tyr Asp Lys Leu Ser Arg Ala Leu 368
 Arg Tyr Tyr Tyr Asp Lys Asn Ile Met Thr Lys Val His Pro Pro Glu 384
 Ser Ser Met Tyr Lys Tyr Pro Ser Asp Leu Pro Tyr Met Ser Ser Tyr 400
 His Gly Lys Arg Tyr Ala Tyr Lys Phe Asp Phe His Gly Ile Ala Gln 416
 Ala Leu Gln Pro His Ala His Pro Gln Lys Met Asn Phe Val Ala Pro 432

His Pro Pro Ala Leu Pro Val Thr Ser Ser Ser Phe Phe Ala Ala Pro 448
Asn Pro Tyr Trp Asn Ser Pro Thr Gly Gly Ile Tyr Pro Asn Thr Arg 464
Leu Pro Ala Ala His Met Pro Ser His Leu Gly Thr Tyr Tyr 478

SEQ ID NO: 5

LENGTH: 23

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION:

ATCTTGATCA CATTATGGCA AGC 23

SEQ ID NO: 6

LENGTH: 25

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION:

CACATTATGG CAAGCACTAT TAAGG 25

SEQ ID NO: 7

LENGTH: 25

TYPE: nucleic acid

CGS 97-01

STANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION:

CACCTAGTAG TAGGTGCCAA GATGG 25